

## AMENDMENTS

In the specification:

On page 46, please delete the paragraph on lines 6-17 and substitute therefor:

Using the ATG identified in the genomic clone as the start codon for translation yielded a novel protein sequence of 1252 amino acids (Figure 2B). Protein domain analysis of the Nope Sequence revealed the presence of a signal peptide, four immunoglobulin (Ig) domains, five fibronectin-type III (FnIII) repeats, a transmembrane domain and a cytoplasmic domain of 274 amino acids (see Figure 1B). Figure ~~[[4C]]~~ 1B shows the domain structure of the Nope protein in comparison to DCC, Punc, and NCAM. Shading indicates similarity among Ig domains. The domain structure indicates that the Nope protein most closely resembles cell adhesion molecules.